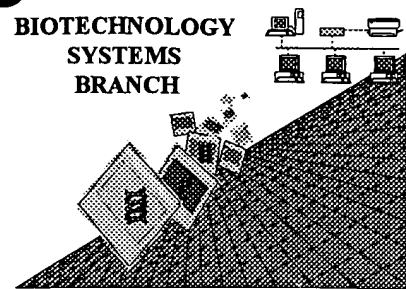


RAW SEQUENCE LISTING

ERROR REPORT

BIOTECHNOLOGY
SYSTEMS
BRANCH



The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) detected errors when processing the following CRF diskette:

Application Serial Number: 09/157,289
Art Unit / Team No. : 01PE
Date Processed by STIC: 9/30/98

THE ATTACHED PRINTOUT EXPLAINS THE ERRORS DETECTED.

PLEASE BE SURE TO FORWARD THIS INFORMATION TO THE APPLICANTS BY EITHER:

- 1) INCLUDING A COPY OF THIS PRINTOUT IN YOUR NEXT COMMUNICATION TO THE APPLICANTS ALONG WITH A NOTICE TO COMPLY or,**
- 2) CALLING APPLICANTS AND FAXING THEM A COPY OF THE PRINTOUT WITH A NOTICE TO COMPLY**

THIS WILL INSURE THAT THE NEXT SUBMISSION RECEIVED FROM THEM WILL BE ERROR FREE.

IF YOU HAVE ANY FURTHER QUESTIONS, PLEASE CALL:

ARTI SHAH 703-308-4212

Law Sequence Listing Error Summary

ERROR DETECTED SUGGESTED CORRECTION

SERIAL NUMBER: 09/152,289

ATTN: NEW RULES CASES: PLEASE DISREGARD ENGLISH "ALPHA" HEADERS, WHICH WERE INSERTED BY PTO SOFTWARE

- 1 **Wrapped Nucleics** The number/text at the end of each line "wrapped" down to the next line.
This may occur if your file was retrieved in a word processor after creating it.
Please adjust your right margin to .3, as this will prevent "wrapping".
- 2 **Wrapped Aminos** The amino acid number/text at the end of each line "wrapped " down to the next line.
This may occur if your file was retrieved in a word processor after creating it.
Please adjust your right margin to .3, as this will prevent "wrapping".
- 3 **Incorrect Line Length** The rules require that a line not exceed 72 characters in length. This includes spaces.
All text must be visible on page.
- 4 **Misaligned Amino Acid Numbering** The numbering under each 5th amino acid is misaligned. This may be caused by the use of tabs between the numbering. It is recommended to delete any tabs and uses spacing between the numbers.
- 5 **Non-ASCII** This file was not saved in ASCII (DOS) text, as required by the Sequence Rules.
Please ensure your subsequent submission is saved in ASCII text so that it can be processed.
- 6 **Variable Length** Sequence(s) contain n's or Xaa's which represented more than one residue.
As per the rules, each n or Xaa can only represent a single residue.
Please present the maximum number of each residue having variable length and indicate in the (ix) features section that some may be missing.
- 7 **Wrong Designation** Sequence(s) contain amino acid or nucleic acid designators which are not standard representations as per the Sequence Rules (Please refer to paragraph 1.822)
- 8 **Skipped Sequences (OLD RULES)** Sequence(s) missing. If intentional, please use the following format for each skipped sequence:
(2) INFORMATION FOR SEQ ID NO:X:
(I) SEQUENCE CHARACTERISTICS:(Do not insert any headings under "SEQUENCE CHARACTERISTICS")
(xI) SEQUENCE DESCRIPTION:SEQ ID NO:X:
This sequence is intentionally skipped

Please also adjust the "(iii) NUMBER OF SEQUENCES:" response to include the skipped sequence(s).
- 9 **Skipped Sequences (NEW RULES)** Sequence(s) missing. If intentional, please use the following format for each skipped sequence.
<210> sequence id number
<400> sequence id number
000
- 10 J **Use of N's or Xaa's (NEW RULES)** Use of N's and/or Xaa's have been detected in the Sequence Listing.
Use of <220> to <223> is MANDATORY if n's or Xaa's are present.
- 11 **Use of <213>Organism (NEW RULES)** Sequence(s) are missing this mandatory field or its response.
- 12 **Use of <220>Feature (NEW RULES)** Sequence(s) are missing the <220>Feature and associated headings.
Use of <220> to <223> is MANDATORY if <213>ORGANISM is "Artificial" or "Unknown"
(See "Federal Register," 6/01/98, Vol. 63, No. 104, pp. 29631-32)
(Sec. 1.823 of new Sequence Rules)
- 13 **Wrong Format** File submitted was in the alphabetical heading format of the Old Sequence Rules. This is invalid since the "Requirements for Patent Applications Containing Nucleotide Sequence and/or Amino Acid Disclosures" Federal Register Notice, Vol. 63, No. 104, June 1, 1998, p. 29620 applies to applications filed on or after July 1, 1998.
AKS-Biotechnology Systems Branch- 7/10/98

RAW SEQUENCE LISTING
PATENT APPLICATION US/09/157,289DATE: 09/30/98
TIME: 11:28:39

Input Set: I157289.RAW

This Raw Listing contains the General Information
Section and up to first 5 pages.

*new format**see pp. 2-4*

Does Not Comply
Corrected Diskette Needed

<110> APPLICANT: ASHKENAZI, AVI J.
BOTSTEIN, DAVID
DODGE, KELLY H.
GURNEY, AUSTIN L.
KIM, KYUNG JIN
LAWRENCE, DAVID A.
PITTI, ROBERT
ROY, MARGARET A.
TUMAS, DANIEL B.
WOOD, WILLIAM I.
GENENTECH INC.
<120> TITLE OF INVENTION: DcR3 Polypeptide, A TNFR Homolog
<130> FILE REFERENCE: 11669.31US03
<140> CURRENT APPLICATION NUMBER: US/09/157,289
<141> CURRENT FILING DATE: 1998-09-18
<150> EARLIER APPLICATION NUMBER: 60/059,288
<151> EARLIER FILING DATE: 1997-09-18
<150> EARLIER APPLICATION NUMBER: 60/094,640
<151> EARLIER FILING DATE: 1998-07-30
<160> NUMBER OF SEQ ID NOS: 16
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<213> ORGANISM: Homo sapiens
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20 25 30
Thr Pro Thr Tyr Pro Trp Arg Asp Ala Glu Thr Gly Glu Arg Leu Val
35 40 45
Cys Ala Gln Cys Pro Pro Gly Thr Phe Val Gln Arg Pro Cys Arg Arg
50 55 60
Asp Ser Pro Thr Thr Cys Gly Pro Cys Pro Pro Arg His Tyr Thr Gln
65 70 75 80
Phe Trp Asn Tyr Leu Glu Arg Cys Arg Tyr Cys Asn Val Leu Cys Gly
85 90 95
Glu Arg Glu Glu Glu Ala Arg Ala Cys His Ala Thr His Asn Arg Ala
100 105 110
Cys Arg Cys Arg Thr Gly Phe Phe Ala His Ala Gly Phe Cys Leu Glu
115 120 125
His Ala Ser Cys Pro Pro Gly Ala Gly Val Ile Ala Pro Gly Thr Pro
130 135 140

PAGE: 2

RAW SEQUENCE LISTING

PATENT APPLICATION US/09/157,289

 DATE: 09/30/98
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 46 145 150 155 160
 47 Ser Ser Ser Ser Ser Glu Gln Cys Gln Pro His Arg Asn Cys Thr Ala
 48 165 170 175
 49 Leu Gly Leu Ala Leu Asn Val Pro Gly Ser Ser Ser His Asp Thr Leu
 50 180 185 190
 51 Cys Thr Ser Cys Thr Gly Phe Pro Leu Ser Thr Arg Val Pro Gly Ala
 52 195 200 205
 53 Glu Glu Cys Glu Arg Ala Val Ile Asp Phe Val Ala Phe Gln Asp Ile
 54 210 215 220
 55 Ser Ile Lys Arg Leu Gln Arg Leu Leu Gln Ala Leu Glu Ala Pro Glu
 56 225 230 235 240
 57 Gly Trp Gly Pro Thr Pro Arg Ala Gly Arg Ala Ala Leu Gln Leu Lys
 58 245 250 255
 59 Leu Arg Arg Arg Leu Thr Glu Leu Leu Gly Ala Gln Asp Gly Ala Leu
 60 260 265 270
 61 Leu Val Arg Leu Leu Gln Ala Leu Arg Val Ala Arg Met Pro Gly Leu
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 64 290 295 300

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66 <211> LENGTH: 1114

67 <212> TYPE: DNA

68 <213> ORGANISM: Homo sapiens

69 <400> SEQUENCE: 2

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 72 aggcctgtcg ctgctgtgcc tgggtgttggc gctgcctgcc ctgctgccgg tggcggtgt 180
 73 acgcggagtg gcagaaacac ccacctaccc ctggcgggac gcagagacag gggagcggct 240
 74 ggtgtgcgcc cagtgcctcc caggcacctt tgtgcagcgg ccgtgccgcc gagacagccc 300
 75 caccagctgt ggcccgtgtc caccgcgcca ctacacgcag ttctggaact acctggagcg 360
 76 ctgccgctac tgcaacgtcc tctgcgggga gcgtgaggag gaggcacggg cttgccacgc 420
 77 caccacaac cgtgcctgcc gctgccgcac cggcttcttc gcgcacgctg gtttctgctt 480
 78 ggagcacgca tcgtgtccac ctggtgccgg cgtgattgcc ccgggcaccc ccagccagaa 540
 79 caccagctgc cagccgtgcc cccaggcac cttctcagcc agcagctcca gctcagagca 600
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 86 gctggagcgg agcgtccgtg agcgttctc ccctgtgcac tgatcctggc cccctcttat 1020
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90 <211> LENGTH: 491

91 <212> TYPE: DNA

92 <213> ORGANISM: Unknown

93 <220> FEATURE:

94 <223> OTHER INFORMATION: Description of Unknown Organism: UNKNOWN

See item 10 on
Even summary sheet

PAGE: 3

RAW SEQUENCE LISTING PATENT APPLICATION US/09/157,289

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See item 10

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98 gggcttgcca cgccaccac aaccgtgcct gccgtgccg caccggcttc ttgcgcacg 180
99 ctggtttctg cttggagcac gcatcgtgtc cacctggtgc cggcgtgatt gccccgggca 240
100 cccccagcca gaacacgcag tgccatagccg tgccccccag gcaccttctc agccagcagc 300
101 tccagctcag agcagtgccg gccccaccgc aactgcacgg ccctgggcct ggccctcaat 360
102 gtgccaggct cttcctccca tgacaccctg tgcaccagct gcaactggctt cccctcagc 420
103 accagggtac caggagctga ggagtgtgag cgtgccgtca tcgactttgt ggctttccag 480
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106 <211> LENGTH: 73
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108 <213> ORGANISM: Unknown
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113 ctacctggag cgc 73
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W--> 123 gngcttgcca cgccaccac aaccgcgcct gcnctgcag caccggnttc ttgcgcacg 180
W--> 124 ctgntttctg cttggagcac gcatcgtgtc cacctggtgn cggcgtgatt gcnccgggca 240
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134 ggaggaggca cgggcttgcc acgccacca caaccgtgcc tgccgtgcc gcaccggctt 120
135 cttgcgcac gctggtttct gcttgagca cgcacgtgtt ccacctggtg ccggcgtgat 180
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142 <223> OTHER INFORMATION: Description of Unknown Organism: UNKNOWN
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item 10

PAGE: 4

RAW SEQUENCE LISTING PATENT APPLICATION US/09/157,289

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item 10

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145      ttgcaccctg agctaggaca ccagttcccc tgaccctgtt cttccctcct ggctgcaggc 120
W--> 146      acccccagcc agaacacgca gnccagccgt gccccccagg caccttctca gccagcagct 180
147      ccagctcaga gcagtgccag ccccaccgca actgcacggc cctgggcctg gccctcaatg 240
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W--> 169      cgctgtgcac cagctgcact ggcttcccc tccagcaccag ggtancagga gctgaggagt 180
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171      <210> SEQ ID NO 10
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173      <212> TYPE: DNA
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182      <211> LENGTH: 21
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184      <213> ORGANISM: Unknown
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186      <223> OTHER INFORMATION: Description of Unknown Organism: UNKNOWN
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192      <213> ORGANISM: Unknown
193      <220> FEATURE:
194      <223> OTHER INFORMATION: Description of Unknown Organism: UNKNOWN

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RAW SEQUENCE LISTING
PATENT APPLICATION US/09/157,289DATE: 09/30/98
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Input Set: I157289.RAW

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209	<220> FEATURE:	
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224	<213> ORGANISM: Unknown	
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228	atcacgccgg caccag	16

Input Set: I157289.RAW

Line	?	Error/Warning	Original Text
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97	W	"N" or "Xaa" used: Feature required	antaactgga gcncctgccgc tactgnaacg tcctctgn
121	W	"N" or "Xaa" used: Feature required	gccgagacag cccacgacg tgtggcccggt gtccaccg
122	W	"N" or "Xaa" used: Feature required	antaactgga gcncctgccgc tactgnaacg tcctctgn
123	W	"N" or "Xaa" used: Feature required	gngcttgcca cgccaccac aaccgcgcct gcngctgc
124	W	"N" or "Xaa" used: Feature required	ctgntttctg cttggagcac gcatcgtgtc cacctggt
136	W	"N" or "Xaa" used: Feature required	tnccccgggc accccagcc a
146	W	"N" or "Xaa" used: Feature required	acccccagcc agaacacgca gnccagccgt gccccca
167	W	"N" or "Xaa" used: Feature required	agcngtgcnc cncaggcacc ttctcagcca gcagttcc
169	W	"N" or "Xaa" used: Feature required	cgctgtgcac cagctgcact ggcttcccc tcagcacc
176	W	"N" or "Xaa" used: Feature required	cttgtccacc tggtgccggc gtgattnccc gggcacc
177	W	"N" or "Xaa" used: Feature required	gccntcccc caggcacctt ctcagccagc agctccag
178	W	"N" or "Xaa" used: Feature required	cgcaactgca acgccctggn ctggccctca atgtgcca